

#### IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Patent Application of: Confirmation No.: 4511

Nordine CHEIKH et al. Art Unit: 1631

Appln. No.: 09/300,482 Examiner: Marjorie A. Moran

Filed: April 28, 1999 Atty. Docket: 16517.216

For: Nucleic Acid Molecules and Other Molecules Associated with the

Phosphogluconate Pathway

### DECLARATION UNDER 37 C.F.R. § 1.132

Assistant Commissioner for Patents Washington, DC 20231

Sir:

BEST AVAILABLE COPY

- I, Nancy Vosnidou, Ph.D., make the following declaration related to matters that are within my personal knowledge:
- 1. I am employed as a Patent Scientist at the Monsanto Company, which wholly owns Monsanto Technology LLC, the assignee of the above referenced patent application.
- 2. I have reviewed the above-captioned application, which was filed April 28, 1999 and assigned Serial No. 09/300,482 (hereinafter the "'482 application"). The '482 application discloses 699 expressed sequence tag sequences (ESTs) derived from corn and soybean having homology to known phosphogluconate pathway enzymes. SEQ ID NO: 1, is identified in the specification as having homology to glucose-6-phosphate-1-dehydrogenase.
- 3. I hereby certify that I conducted a computer search of an identical copy of the nucleotide database posted by the National Center for Biotechnology Information (NCBI) contained on a

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Page 2

Monsanto local server, on March 30, 2006. The computer search was conducted using a BLASTN query of SEQ ID NO: 1 using the default parameters. The search was further conducted without identifying open reading frames (ORFs) within the sequence prior to the

4. I hereby certify that a copy of an alignment between SEQ ID NO: 1 and the top matches resulting from the computer search is attached hereto as appendix A. The alignment shows that SEQ ID NO: 1 has 95% identity to GenBank Accession No. 60672339. This GenBank accession is identified as similar to glucose-6-phosphate dehydrogenase.

I further declare that all statements made herein of my own knowledge are true and that all statements made on information and belief are believed to be true; and further that these statements were made with the knowledge that willful false statements and the like so made are punishable by fine or imprisonment, or both, under Section 1001 of Title 18 of the United States Code, and that such willful false statements may jeopardize the validity of the application or any patent issued thereon.

Nancy Vosnidou, Ph.D.

Executed on <u>24 April</u>, 2006.

search.



<HR>
<PRE>
<B>BLASTN 2.2.9 [May-01-2004]</B>
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Reference: Altschul, Stephen F., Thomas L. Madden, Alejandro A. Schaffer, Jinghui Zhang, Zheng Zhang, Webb Miller, and David J. Lipman (1997), "Gapped BLAST and PSI-BLAST: a new generation of protein database search programs", Nucleic Acids Res. 25:3389-3402.

Query= 15365\_SEQID1 (233 letters)

Searching......done

Sequences producing significant alignments:	Score (bits)	E Value	
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                  Sbjct: 604 ttaaacaaggcaatctctgagtatgaaacatcaaaaaacaatgaatcaggaagctatcgc 663
Query: 122 agattatttttatttggcattgcctccatcagtctacccatcagtatgcgagatgataaga 181
                   Sbjct: 664 agattattttatttggcattgcctccatcagtctacccatcagtatgcaaaatgataaga 723
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l">Sequence Details</a> <a
href="http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?cmd=Search&db=Nucleotide&ter
\verb|m=36019255">NCBI</a> description="SCMCSD1059C02.g SD1 Saccharum officinarum officinaru
                  cDNA clone SCMCSD1059C02 5'." library=NA
                   species="Saccharum officinarum" source="genbank_est"
                  version=NA type=cDNA
                 Length = 711
  Score = 311 \text{ bits } (157), \text{ Expect = } 7e-82
  Identities = 211/229 (92%)
  Strand = Plus / Plus
Query: 2
                   ttttttgcagttagtagaatatgttagtggctcctatgatagggtggaaggatttgagtta 61
                   Sbjct: 59 tttttgcagctagttaaatatgttagtggctcctatgatagtggggaaggatttgagtta 118
Query: 62
                  ttgaatgaggcaatctctgagtatgagacttcagaaaacaatgactcgggaagctaccgc 121
                   Sbjct: 119 ttaaacaaggcactctctgagtatgaaacatcaaaaaacaatgaatcgggaagctatcgc 178
Query: 122 agattattttatttggcattgcctccatcagtctacccatcagtatgcgagatgataaga 181
                   Sbjct: 179 agattattttatttggcattgcctccatcagtctacccatcagtatgcaaaatgataaga 238
Query: 182 tcatattgcatgagtccatcttcacacacggttggacaagggttattg 230
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href="../../GenesisDataRequest/GenesisDataRequest?QUERY=%7B%5B%28Seq_id%2Fid_val
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l">Sequence Details</a> <a
href="http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?cmd=Search&db=Nucleotide&ter
m=60672339">NCBI</a> description="EST5840 Zea mays embryo sac cDNA library Zea
         mays cDNA clone ES12956 5' similar to
         glucose-6-phosphate dehydrogenase." library=NA
         species="Zea mays" source="genbank_est" version=NA
         type=cDNA
        Length = 621
Score = 309 \text{ bits } (156), Expect = 3e-81
 Identities = 180/188 (95%)
 Strand = Plus / Plus
Query: 46 ggaaggatttgagttattgaatgaggcaatctctgagtatgagacttcagaaaacaatga 105
          ggaaggatttgagttattaaataaggcaatctctgagtatgaaacttcaaaaaacaatga 64
Sbjct: 5
Query: 106 ctcgggaagctaccgcagattattttatttggcattgcctccatcagtctacccatcagt 165
          Sbjct: 65 ctcgggaagctaccacagattattttatttggcattgcctccatcagtctacccatcagt 124
Query: 166 atgcgagatgataagatcatattgcatgagtccatcttcacacaccggttggacaagggt 225
          Sbjct: 125 atgcaaaatgataagatcatattgcatgagtccatcttcacacactggttggacaagggt 184
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          1111111
Sbjct: 185 tattgttg 192
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l">Sequence Details</a> <a
href="http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?cmd=Search&db=Nucleotide&ter
m=62550618">NCBI</a> description="EST7250 Zea mays embryo sac cDNA library Zea
          mays cDNA clone ES14608 5' similar to
          glucose-6-phosphate dehydrogenase." library=NA
          species="Zea mays" source="genbank_est" version=NA
          type=cDNA
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 Score = 309 \text{ bits (156)}, Expect = 3e-81
 Identities = 180/188 (95%)
 Strand = Plus / Plus
Query: 46 ggaaggatttgagttattgaatgaggcaatctctgagtatgagacttcagaaaacaatga 105
          Sbjct: 5
          ggaaggatttgagttattaaataaggcaatctctgagtatgaaacttcaaaaaacaatga 64
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Query: 166 atgcgagatgataagatcatattgcatgagtccatcttcacacaccggttggacaagggt 225
         Sbjct: 125 atgcaaaatgataagatcatattgcatgagtccatcttcacacactggttggacaagggt 184
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         1111111
Sbjct: 185 tattgttg 192
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href="../../GenesisDataRequest/GenesisDataRequest?QUERY=%7B%5B%28Seq_id%2Fid_val
ue%3D%2735138042%27%29%5D%7D&DATABASE=GCGEST&LENGTH=1&VIEW=SequenceDetailView.xs
l">Sequence Details</a> <a
href="http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?cmd=Search&db=Nucleotide&ter
m=35138042">NCBI</a> description="SCCCST3C01H12.g ST3 Saccharum officinarum
         cDNA clone SCCCST3C01H12 5'." library=NA
         species="Saccharum officinarum" source="genbank_est"
         version=NA type=cDNA
        Length = 689
 Score = 303 \text{ bits } (153), \text{ Expect = } 2e-79
 Identities = 210/229 (91%)
 Strand = Plus / Plus
Query: 2
         tttttgcagttagtagaatatgttagtggctcctatgatagggtggaaggatttgagtta 61
         Sbjct: 405 tttttgcagctagttaaatatgttagtggctcctatgatagtggggaaggatttgagttg 464
Query: 62
         ttgaatgaggcaatctctgagtatgagacttcagaaaacaatgactcgggaagctaccgc 121
         Sbjct: 465 ttaaacaaggcaatctctgagtatgaaacatcaaaaaacaatgaatcaggaagctatcgc 524
Query: 122 agattattttatttggcattgcctccatcagtctacccatcagtatgcgagatgataaga 181
          Sbjct: 525 agattattttatttggcattgcctccatcagtctacccatcagtatgcaaaatgataaga 584
Query: 182 tcatattgcatgagtccatcttcacacacggttggacaagggttattg 230
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المرايد بالمراجع المتعلقة والمتعلقة والمتعلق والمتعلقة والمتعلقة والمتعلقة والمتعلقة والمتعلقة والمتعلقة و

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l">Sequence Details</a> <a
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m=57813598">NCBI</a> description="GABR1_16_C06.g1_A002 GA- or
         brassinolide-treated seedlings Sorghum bicolor cDNA
         clone GABR1_16_C06_A002 5'." library=NA species="Sorghum
         bicolor" source="genbank_est" version=NA type=cDNA
        Length = 786
 Score = 301 \text{ bits } (152), Expect = 7e-79
 Identities = 212/232 (91%)
 Strand = Plus / Plus
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Query: 2
         Sbjct: 545 tttttgcagctagttaaatatgttagtggctcctatgatagtggggaaggatttgagtta 604
Query: 62 ttgaatgaggcaatctctgagtatgagacttcagaaaacaatgactcgggaagctaccgc 121
          Sbjct: 605 ttaaacaaggcaatctccgagtatgaaacatcaaagaacaatgaatcaggaagctatcgc 664
Query: 122 agattattttatttggcattgcctccatcagtctacccatcagtatgcgagatgataaga 181
          Sbjct: 665 agattattttatttggcattgcctccatcagtctacccatcagtatgcaaaatgataaga 724
Query: 182 tcatattgcatgagtccatcttcacacacggttggacaagggttattgttg 233
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          cDNA clone SCACCL6008B07 5'." library=NA
          species="Saccharum officinarum" source="genbank_est"
          version=NA type=cDNA
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 Score = 293 bits (148), Expect = 2e-76
 Identities = 211/232 (90%)
 Strand = Plus / Plus
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Query: 2
          Sbjct: 243 tttttgcagctagttaaatatgttagtggctcctatgatagtggggaaggatttgagtca 302
Query: 62 ttgaatgaggcaatctctgagtatgagacttcagaaaacaatgactcgggaagctaccgc 121
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Sbjct: 303 ttaaacaaggcaatctctgagtatgaaacatcaaaaaacaatgaatcaagaagctatcgc 362
Query: 122 agattattttatttggcattgcctccatcagtctacccatcagtatgcgagatgataaga 181
         Sbjct: 363 agattattttatttggcattgcctccatcagtctacccatcagtatgcaaaatgataaga 422
Query: 182 tcatattgcatgagtccatcttcacacaccggttggacaagggttattgttg 233
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l">Seguence Details</a> <a
href="http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?cmd=Search&db=Nucleotide&ter
m=45981870">NCBI</a> description="WOUND1_18_A03.g1_A002 Wounded leaves Sorghum
         bicolor cDNA clone WOUND1_18_A03_A002 5'." library=NA
         species="Sorghum bicolor" source="genbank_est"
         version=NA type=cDNA
        Length = 728
 Score = 287 bits (145), Expect = 1e-74
 Identities = 212/233 (90%), Gaps = 1/233 (0%)
 Strand = Plus / Plus
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          Sbjct: 496 tttttgcagctagttaaatatgttagtggctcctatgatagtggggaaggatttgagtta 555
Query: 62 ttgaatgaggcaatctctgagtatgagacttcagaaaa-caatgactcgggaagctaccg 120
          Sbjct: 556 ttaaacaaggcaatctctgagtatgaaacatcaaaaaacaatgaatcaggaagctatcg 615
Query: 121 cagattattttatttggcattgcctccatcagtctacccatcagtatgcgagatgataag 180
          Sbjct: 616 cagatgattttatttggcattgcctccgtcagtctacccatcagtatgcaaaatgataag 675
Query: 181 atcatattgcatgagtccatcttcacacaccggttggacaagggttattgttg 233
          Sbjct: 676 atcgtattgcatgaatccatcttcacacaccggttggacaagggttattgttg 728
<input type=checkbox name="seqID" value="45997222:genbank_est:" checked>&gt;<a</pre>
href="sequenceViewer?seqID=45997222&source=genbank_est">45997222</a> <a
href="../../GenesisDataRequest/GenesisDataRequest?QUERY=%7B%5B%28Seq_id%2Fid_val
ue%3D%2745997222%27%29%5D%7D&DATABASE=GCGEST&LENGTH=1&VIEW=SequenceDetailView.xs
l">Sequence Details</a> <a
href="http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?cmd=Search&db=Nucleotide&ter
m=45997222">NCBI</a> description="WOUND1_84_F10.g1_A002 Wounded leaves Sorghum
```

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```
version=NA type=cDNA
        Length = 759
Score = 270 bits (136), Expect = 2e-69
Identities = 196/216 (90%)
Strand = Plus / Plus
Query: 2
         tttttgcagttagtagaatatgttagtggctcctatgatagggtggaaggatttgagtta 61
         Sbjct: 544 tttttgcagctagttaaatatgttagtggctcctatgatagtggggaaggatttgagtta 603
Query: 62 ttgaatgaggcaatctctgagtatgagacttcagaaaacaatgactcgggaagctaccgc 121
         Sbjct: 604 ttaaacaaggcaatctctgagtatgaaacatcaaaaaacaatgaatcacgaagctatcgc 663
Query: 122 agattattttatttggcattgcctccatcagtctacccatcagtatgcgagatgataaga 181
         Sbjct: 664 agattattttatttggcattgcctccatcagtctacccatcagtatgcaaaattataaga 723
Query: 182 tcatattgcatgagtccatcttcacacaccggttgg 217
         Sbjct: 724 tcgtattgcatgaatccatcttcacacaccggttgg 759
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href="../../GenesisDataRequest/GenesisDataRequest?QUERY=%7B%5B%28Seq_id%2Fid_val
ue%3D%2734443163%27%29%5D%7D&DATABASE=GCGEST&LENGTH=1&VIEW=SequenceDetailView.xs
l">Sequence Details</a> <a
href="http://www.ncbi.nlm.nih.gov/entrez/query.fcqi?cmd=Search&db=Nucleotide&ter
m=34443163">NCBI</a> description="PH1_28_C06.g1_A002 Phosphorous-deficient
         seedlings Sorghum bicolor cDNA clone PH1_28_C06_A002
         5'." library=NA species="Sorghum bicolor"
         source="genbank_est" version=NA type=cDNA
        Length = 795
 Score = 264 bits (133), Expect = 1e-67
 Identities = 205/227 (90%), Gaps = 4/227 (1%)
 Strand = Plus / Plus
Ouerv: 2
         ttttttgcagttagtagaatatgttagtggctcctatgata---gggtggaaggatttgag 58
         Sbjct: 570 tttttgcagctagttaaatatgttagtggctcctat-ataagtgggggaaaggatttgag 628
Query: 59 ttattgaatgaggcaatctctgagtatgagacttcagaaaacaatgactcgggaagctac 118
         Sbjct: 629 ttattaaacaaggcaatctctgagtatgaaacatcaaaaaacaatgaatcaggaagctat 688
```

American Ame

bicolor cDNA clone WOUND1\_84\_F10\_A002 5'." library=NA

species="Sorghum bicolor" source="genbank\_est"

```
Query: 119 cgcagattattttatttggcattgcctccatcagtctacccatcagtatgcgagatgata 178
          Sbjct: 689 cgcagattattttatttggcattgcctccatcagtctacccatcagtatgcaaaatgata 748
Query: 179 agatcatattgcatgagtccatcttcacacaccggttggacaagggt 225
          Sbjct: 749 agatcgtattgcatgaatccatcttcacacaccggttggacaagggt 795
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ue%3D%2757812762%27%29%5D%7D&DATABASE=GCGEST&LENGTH=1&VIEW=SequenceDetailView.xs
l">Sequence Details</a> <a
href="http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?cmd=Search&db=Nucleotide&ter
m=57812762">NCBI</a> description="GABR1_11_E09.g1_A002 GA- or
         brassinolide-treated seedlings Sorghum bicolor cDNA
         clone GABR1_11_E09_A002 5'." library=NA species="Sorghum
         bicolor" source="genbank_est" version=NA type=cDNA
         Length = 746
 Score = 258 bits (130), Expect = 9e-66
 Identities = 184/202 (91%)
 Strand = Plus / Plus
Query: 2
          tttttgcagttagtagaatatgttagtggctcctatgatagggtggaaggatttgagtta 61
          Sbjct: 545 ttttttgcagctagttaaatatgttagtggctcctatgatagtggggaaggatttgagtta 604
Query: 62 ttgaatgaggcaatctctgagtatgagacttcagaaaacaatgactcgggaagctaccgc 121
          Sbjct: 605 ttaaacaaggcaatctctgagtatgaaacatcaaaaaacaatgaatcaggaagctatcgc 664
Query: 122 agattattttatttggcattgcctccatcagtctacccatcagtatgcgagatgataaga 181
          Sbjct: 665 agattattttatttggcattgcctccatcagtctacccatcagtatgcaaaatgataaga 724
Query: 182 tcatattgcatgagtccatctt 203
          11 11111111
Sbjct: 725 tcgtattgcatgaatccatctt 746
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l">Sequence Details</a> <a
href="http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?cmd=Search&db=Nucleotide&ter
m=35138932">NCBI</a> description="SCQGAM2107H03.g AM2 Saccharum officinarum
          cDNA clone SCQGAM2107H03 5'." library=NA
          species="Saccharum officinarum" source="genbank_est"
          version=NA type=cDNA
```

Score = 256 bits (129), Expect = 4e-65 Identities = 156/165 (94%)Strand = Plus / Plus Query: 69 aggcaatctctgagtatgagacttcagaaaacaatgactcgggaagctaccgcagattat 128 Sbjct: 3 aggcaatctctgagtatgaaacatcaaaaaacaatgaatcgggaagctatcgcagattat 62 Query: 129 tttatttggcattgcctccatcagtctacccatcagtatgcgagatgataagatcatatt 188 Sbjct: 63 thtatttggcattgcctccatcagtctacccatcagtatgcaaaatgataagatcgtatt 122 Query: 189 gcatgagtccatcttcacacaccggttggacaagggttattgttg 233 Sbjct: 123 gcatgaatccatcttcacacacggttggacaagggttattgttg 167 <input type=checkbox name="seqID" value="35278891:genbank\_est:" checked>&gt;<a</pre> href="sequenceViewer?seqID=35278891&source=genbank\_est">35278891</a> <a href="../../GenesisDataRequest/GenesisDataRequest?QUERY=%7B%5B%28Seq\_id%2Fid\_val ue%3D%2735278891%27%29%5D%7D&DATABASE=GCGEST&LENGTH=1&VIEW=SequenceDetailView.xs l">Sequence Details</a> <a href="http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?cmd=Search&db=Nucleotide&ter m=35278891">NCBI</a> description="SCEZFL4039C11.g FL4 Saccharum officinarum cDNA clone SCEZFL4039C11 5'." library=NA species="Saccharum officinarum" source="genbank\_est" version=NA type=cDNA Length = 768Score = 232 bits (117), Expect = 5e-58Identities = 205/233 (87%), Gaps = 1/233 (0%) Strand = Plus / Plus tttttgcagttagtagaatatgttagtggctcctatgatagggtggaaggatttgagtta 61 Query: 2 Sbjct: 465 tttttgcagctagttaaatatgttagtggctcctatgatagtggggaaggatttgagtta 524 Ouery: 62 ttgaatgaggcaatctctgagtatgagacttcagaaaacaatgactcgggaagctaccgc 121 Sbjct: 525 ttaaacaaggcaatctctgagtatgaaacatcaaaaaacaatgaatcgggaagctatcgc 584 Query: 122 agattattttatttggcattgcctccatcagtctacccatcagtatgcgagatgataaga 181

Sbjct: 585 agattattttattttgcatttgcttcatcagtctacccatcagtatgcaaaatgattaga 644

Query: 182 tcatattgcatgagtccatcttcacacaccggttgga-caagggttattgttg 233

Sbjct: 645 tcggattgcatgaaatcatcttcacacacccgttggaccaaggtttattgttg 697

Length = 658

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l">Sequence Details</a> <a
href="http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?cmd=Search&db=Nucleotide&ter
m=86469135">NCBI</a> description="ZM_BFb0246E12.r ZM_BFb Zea mays cDNA 5'."
         library=NA species="Zea mays" source="genbank_est"
         version=NA type=cDNA
        Length = 611
 Score = 226 bits (114), Expect = 3e-56
 Identities = 129/134 (96%)
 Strand = Plus / Plus
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          Sbjct: 478 tttttgcagttagtaaaatatgttagtggctcctatgatagggtggaaggatttgagtta 537
Query: 62 ttgaatgaggcaatctctgagtatgagacttcagaaaacaatgactcgggaagctaccgc 121
          Sbjct: 538 ttaaataaggcaatctctgagtatgaaacttcaaaaaacaatgactcgggaagctaccgc 597
Query: 122 agattattttattt 135
          Sbjct: 598 agattattttattt 611
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l">Sequence Details</a> <a
href="http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?cmd=Search&db=Nucleotide&ter
m=49455184">NCBI</a> description="MJR031G10 Methyl jasmonate-treated root
          Lambda ZIPLOX Library (MJR) Saccharum sp. cDNA clone
          MJR062E04 5' similar to glucose-6-phosphate
          dehydrogenase." library=NA species="Saccharum sp."
          source="genbank_est" version=NA type=cDNA
         Length = 376
 Score = 212 bits (107), Expect = 5e-52
 Identities = 143/156 (91%)
 Strand = Plus / Plus
Query: 78 ctgagtatgagacttcagaaaacaatgactcgggaagctaccgcagattattttatttgg 137
```

Query: 138 cattgcctccatcagtctacccatcagtatgcgagatgataagatcatattgcatgagtc 197

```
Sbjct: 78 cattgcctccatcagtctacccatcagtatgcaaaatgataagatcgtattgcatgaatc 137
Query: 198 catcttcacacaccggttggacaagggttattgttg 233
          Sbjct: 138 catcttcacacaccggttggacaagggttattgttg 173
 Database: Predefined collection GenBankCDNA: -source
"DDBJ_EST, DDBJ_NUC, EMBL_EST, EMBL_NUC, GENBANK_EST, GENBANK_NUC, OTHER_EST, OTHER_NUC
, PDB_EST, PDB_NUC, UNKNOWN_EST, UNKNOWN_NUC" -type "CDNA" -maxLength 250000 -chunk
    Posted date: Mar 28, 2006 2:13 PM
  Number of letters in database: 937,495,304
 Number of sequences in database: 629,443
 Database: Predefined collection GenBankCDNA: -source
"DDBJ_EST, DDBJ_NUC, EMBL_EST, EMBL_NUC, GENBANK_EST, GENBANK_NUC, OTHER_EST, OTHER_NUC
, PDB_EST, PDB_NUC, UNKNOWN_EST, UNKNOWN_NUC" -type "CDNA" -maxLength 250000 -chunk
    Posted date: Mar 28, 2006 2:16 PM
  Number of letters in database: 923,029,544
 Number of sequences in database: 675,893
  Database: Predefined collection GenBankCDNA: -source
"DDBJ_EST,DDBJ_NUC,EMBL_EST,EMBL_NUC,GENBANK_EST,GENBANK_NUC,OTHER_EST,OTHER_NUC
, PDB_EST, PDB_NUC, UNKNOWN_EST, UNKNOWN_NUC" -type "CDNA" -maxLength 250000 -chunk
    Posted date: Mar 28, 2006 2:19 PM
  Number of letters in database: 918,994,864
  Number of sequences in database: 717,011
  Database: Predefined collection GenBankCDNA: -source
"DDBJ_EST,DDBJ_NUC,EMBL_EST,EMBL_NUC,GENBANK_EST,GENBANK_NUC,OTHER_EST,OTHER_NUC
, PDB_EST, PDB_NUC, UNKNOWN_EST, UNKNOWN_NUC" -type "CDNA" -maxLength 250000 -chunk
    Posted date: Mar 28, 2006 2:22 PM
  Number of letters in database: 872,212,624
  Number of sequences in database: 890,994
  Database: Predefined collection GenBankCDNA: -source
"DDBJ_EST, DDBJ_NUC, EMBL_EST, EMBL_NUC, GENBANK_EST, GENBANK_NUC, OTHER_EST, OTHER_NUC
, PDB_EST, PDB_NUC, UNKNOWN_EST, UNKNOWN_NUC" -type "CDNA" -maxLength 250000 -chunk
    Posted date: Mar 28, 2006 2:25 PM
  Number of letters in database: 727,928,668
  Number of sequences in database: 1,612,608
  Database: Predefined collection GenBankCDNA: -source
"DDBJ_EST,DDBJ_NUC,EMBL_EST,EMBL_NUC,GENBANK_EST,GENBANK_NUC,OTHER_EST,OTHER_NUC
, PDB_EST, PDB_NUC, UNKNOWN_EST, UNKNOWN_NUC" -type "CDNA" -maxLength 250000 -chunk
    Posted date: Mar 28, 2006 2:28 PM
  Number of letters in database: 770.454,229
  Number of sequences in database: 1,841,517
  Database: Predefined collection GenBankCDNA: -source
"DDBJ_EST, DDBJ_NUC, EMBL_EST, EMBL_NUC, GENBANK_EST, GENBANK_NUC, OTHER_EST, OTHER_NUC
, PDB_EST, PDB_NUC, UNKNOWN_EST, UNKNOWN_NUC" -type "CDNA" -maxLength 250000 -chunk
    Posted date: Mar 28, 2006 2:31 PM
  Number of letters in database: 751,537,354
  Number of sequences in database: 1,649,902
```

Database: Predefined collection GenBankCDNA: -source "DDBJ\_EST, DDBJ\_NUC, EMBL\_EST, EMBL\_NUC, GENBANK\_EST, GENBANK\_NUC, OTHER\_EST, OTHER\_NUC , PDB\_EST, PDB\_NUC, UNKNOWN\_EST, UNKNOWN\_NUC" -type "CDNA" -maxLength 250000 -chunk Posted date: Mar 28, 2006 2:34 PM Number of letters in database: 804,957,458 Number of sequences in database: 1,377,732 Database: Predefined collection GenBankCDNA: -source "DDBJ\_EST, DDBJ\_NUC, EMBL\_EST, EMBL\_NUC, GENBANK\_EST, GENBANK\_NUC, OTHER\_EST, OTHER\_NUC , PDB\_EST, PDB\_NUC, UNKNOWN\_EST, UNKNOWN\_NUC" -type "CDNA" -maxLength 250000 -chunk Posted date: Mar 28, 2006 2:39 PM Number of letters in database: 674,836,374 Number of sequences in database: 1,794,151 Database: Predefined collection GenBankCDNA: -source "DDBJ\_EST, DDBJ\_NUC, EMBL\_EST, EMBL\_NUC, GENBANK\_EST, GENBANK\_NUC, OTHER\_EST, OTHER\_NUC , PDB\_EST, PDB\_NUC, UNKNOWN\_EST, UNKNOWN\_NUC" -type "CDNA" -maxLength 250000 -chunk Posted date: Mar 28, 2006 2:42 PM Number of letters in database: 771,112,785 Number of sequences in database: 1,417,437 Database: Predefined collection GenBankCDNA: -source "DDBJ\_EST, DDBJ\_NUC, EMBL\_EST, EMBL\_NUC, GENBANK\_EST, GENBANK\_NUC, OTHER\_EST, OTHER\_NUC , PDB\_EST, PDB\_NUC, UNKNOWN\_EST, UNKNOWN\_NUC" -type "CDNA" -maxLength 250000 -chunk Posted date: Mar 28, 2006 2:45 PM Number of letters in database: 778,599,445 Number of sequences in database: 1,379,262 Database: Predefined collection GenBankCDNA: -source "DDBJ\_EST, DDBJ\_NUC, EMBL\_EST, EMBL\_NUC, GENBANK\_EST, GENBANK\_NUC, OTHER\_EST, OTHER\_NUC , PDB\_EST, PDB\_NUC, UNKNOWN\_EST, UNKNOWN\_NUC" -type "CDNA" -maxLength 250000 -chunk Posted date: Mar 28, 2006 2:49 PM Number of letters in database: 795,162,534 Number of sequences in database: 1,358,269 Database: Predefined collection GenBankCDNA: -source "DDBJ\_EST, DDBJ\_NUC, EMBL\_EST, EMBL\_NUC, GENBANK\_EST, GENBANK\_NUC, OTHER\_EST, OTHER\_NUC , PDB\_EST, PDB\_NUC, UNKNOWN\_EST, UNKNOWN\_NUC" -type "CDNA" -maxLength 250000 -chunk Posted date: Mar 28, 2006 2:52 PM Number of letters in database: 795,846,631 Number of sequences in database: 1,349,677 Database: Predefined collection GenBankCDNA: -source "DDBJ\_EST,DDBJ\_NUC,EMBL\_EST,EMBL\_NUC,GENBANK\_EST,GENBANK\_NUC,OTHER\_EST,OTHER\_NUC , PDB\_EST, PDB\_NUC, UNKNOWN\_EST, UNKNOWN\_NUC" -type "CDNA" -maxLength 250000 -chunk Posted date: Mar 28, 2006 2:56 PM Number of letters in database: 758,471,273 Number of sequences in database: 1,493,133 Database: Predefined collection GenBankCDNA: -source "DDBJ\_EST, DDBJ\_NUC, EMBL\_EST, EMBL\_NUC, GENBANK\_EST, GENBANK\_NUC, OTHER\_EST, OTHER\_NUC , PDB\_EST, PDB\_NUC, UNKNOWN\_EST, UNKNOWN\_NUC" -type "CDNA" -maxLength 250000 -chunk Posted date: Mar 28, 2006 2:59 PM

Number of letters in database: 799,469,308 Number of sequences in database: 1,318,767

Database: Predefined collection GenBankCDNA: -source "DDBJ\_EST, DDBJ\_NUC, EMBL\_EST, EMBL\_NUC, GENBANK\_EST, GENBANK\_NUC, OTHER\_EST, OTHER\_NUC , PDB\_EST, PDB\_NUC, UNKNOWN\_EST, UNKNOWN\_NUC" -type "CDNA" -maxLength 250000 -chunk Posted date: Mar 28, 2006 3:02 PM Number of letters in database: 807,279,523 Number of sequences in database: 1,289,115 Database: Predefined collection GenBankCDNA: -source "DDBJ\_EST, DDBJ\_NUC, EMBL\_EST, EMBL\_NUC, GENBANK\_EST, GENBANK\_NUC, OTHER\_EST, OTHER\_NUC , PDB\_EST, PDB\_NUC, UNKNOWN\_EST, UNKNOWN\_NUC" -type "CDNA" -maxLength 250000 -chunk Posted date: Mar 28, 2006 3:06 PM Number of letters in database: 797,258,625 Number of sequences in database: 1,365,557 Database: Predefined collection GenBankCDNA: -source "DDBJ\_EST, DDBJ\_NUC, EMBL\_EST, EMBL\_NUC, GENBANK\_EST, GENBANK\_NUC, OTHER\_EST, OTHER\_NUC , PDB\_EST, PDB\_NUC, UNKNOWN\_EST, UNKNOWN\_NUC" -type "CDNA" -maxLength 250000 -chunk Posted date: Mar 28, 2006 3:11 PM Number of letters in database: 785,281,935 Number of sequences in database: 1,395,907 Database: Predefined collection GenBankCDNA: -source "DDBJ\_EST, DDBJ\_NUC, EMBL\_EST, EMBL\_NUC, GENBANK\_EST, GENBANK\_NUC, OTHER\_EST, OTHER\_NUC , PDB\_EST, PDB\_NUC, UNKNOWN\_EST, UNKNOWN\_NUC" -type "CDNA" -maxLength 250000 -chunk Posted date: Mar 28, 2006 3:15 PM Number of letters in database: 793,930,244 Number of sequences in database: 1,317,716 Database: Predefined collection GenBankCDNA: -source "DDBJ\_EST, DDBJ\_NUC, EMBL\_EST, EMBL\_NUC, GENBANK\_EST, GENBANK\_NUC, OTHER\_EST, OTHER\_NUC , PDB\_EST, PDB\_NUC, UNKNOWN\_EST, UNKNOWN\_NUC" -type "CDNA" -maxLength 250000 -chunk Posted date: Mar 28, 2006 3:19 PM Number of letters in database: 782,893,843 Number of sequences in database: 1,386,920 Database: Predefined collection GenBankCDNA: -source "DDBJ\_EST, DDBJ\_NUC, EMBL\_EST, EMBL\_NUC, GENBANK\_EST, GENBANK\_NUC, OTHER\_EST, OTHER\_NUC , PDB\_EST, PDB\_NUC, UNKNOWN\_EST, UNKNOWN\_NUC" -type "CDNA" -maxLength 250000 -chunk Posted date: Mar 28, 2006 3:22 PM Number of letters in database: 781,959,849 Number of sequences in database: 1,431,387 Database: Predefined collection GenBankCDNA: -source "DDBJ\_EST, DDBJ\_NUC, EMBL\_EST, EMBL\_NUC, GENBANK\_EST, GENBANK\_NUC, OTHER\_EST, OTHER\_NUC , PDB\_EST, PDB\_NUC, UNKNOWN\_EST, UNKNOWN\_NUC" -type "CDNA" -maxLength 250000 -chunk Posted date: Mar 28, 2006 3:25 PM Number of letters in database: 801,325,451 Number of sequences in database: 1,305,614 Database: Predefined collection GenBankCDNA: -source "DDBJ\_EST, DDBJ\_NUC, EMBL\_EST, EMBL\_NUC, GENBANK\_EST, GENBANK\_NUC, OTHER\_EST, OTHER\_NUC , PDB\_EST, PDB\_NUC, UNKNOWN\_EST, UNKNOWN\_NUC" -type "CDNA" -maxLength 250000 -chunk Posted date: Mar 28, 2006 3:28 PM Number of letters in database: 810,443,508 Number of sequences in database: 1,256,195

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Database: Predefined collection GenBankCDNA: -source \*DDBJ\_EST, DDBJ\_NUC, EMBL\_EST, EMBL\_NUC, GENBANK\_EST, GENBANK\_NUC, OTHER\_EST, OTHER\_NUC , PDB\_EST, PDB\_NUC, UNKNOWN\_EST, UNKNOWN\_NUC" -type "CDNA" -maxLength 250000 -chunk Posted date: Mar 28, 2006 3:31 PM Number of letters in database: 791,952,159 Number of sequences in database: 1,257,465 Database: Predefined collection GenBankCDNA: -source "DDBJ\_EST, DDBJ\_NUC, EMBL\_EST, EMBL\_NUC, GENBANK\_EST, GENBANK\_NUC, OTHER\_EST, OTHER\_NUC , PDB\_EST, PDB\_NUC, UNKNOWN\_EST, UNKNOWN\_NUC" -type "CDNA" -maxLength 250000 -chunk Posted date: Mar 28, 2006 3:34 PM Number of letters in database: 814,678,444 Number of sequences in database: 1,182,353 Database: Predefined collection GenBankCDNA: -source "DDBJ\_EST, DDBJ\_NUC, EMBL\_EST, EMBL\_NUC, GENBANK\_EST, GENBANK\_NUC, OTHER\_EST, OTHER\_NUC , PDB\_EST, PDB\_NUC, UNKNOWN\_EST, UNKNOWN\_NUC" -type "CDNA" -maxLength 250000 -chunk Posted date: Mar 28, 2006 3:37 PM Number of letters in database: 812,835,524 Number of sequences in database: 1,237,994 Database: Predefined collection GenBankCDNA: -source "DDBJ\_EST, DDBJ\_NUC, EMBL\_EST, EMBL\_NUC, GENBANK\_EST, GENBANK\_NUC, OTHER\_EST, OTHER\_NUC , PDB\_EST, PDB\_NUC, UNKNOWN\_EST, UNKNOWN\_NUC" -type "CDNA" -maxLength 250000 -chunk Posted date: Mar 28, 2006 3:40 PM Number of letters in database: 818,500,554 Number of sequences in database: 1,353,259 Database: Predefined collection GenBankCDNA: -source "DDBJ\_EST, DDBJ\_NUC, EMBL\_EST, EMBL\_NUC, GENBANK\_EST, GENBANK\_NUC, OTHER\_EST, OTHER\_NUC , PDB\_EST, PDB\_NUC, UNKNOWN\_EST, UNKNOWN\_NUC" -type "CDNA" -maxLength 250000 -chunk Posted date: Mar 28, 2006 3:44 PM Number of letters in database: 792,475,637 Number of sequences in database: 1,458,981 Database: Predefined collection GenBankCDNA: -source "DDBJ\_EST,DDBJ\_NUC,EMBL\_EST,EMBL\_NUC,GENBANK\_EST,GENBANK\_NUC,OTHER\_EST,OTHER\_NUC , PDB\_EST, PDB\_NUC, UNKNOWN\_EST, UNKNOWN\_NUC" -type "CDNA" -maxLength 250000 -chunk Posted date: Mar 28, 2006 3:47 PM Number of letters in database: 741,997,156 Number of sequences in database: 1,382,357 Database: Predefined collection GenBankCDNA: -source "DDBJ\_EST, DDBJ\_NUC, EMBL\_EST, EMBL\_NUC, GENBANK\_EST, GENBANK\_NUC, OTHER\_EST, OTHER\_NUC , PDB\_EST, PDB\_NUC, UNKNOWN\_EST, UNKNOWN\_NUC" -type "CDNA" -maxLength 250000 -chunk Posted date: Mar 28, 2006 3:49 PM Number of letters in database: 253,846,446 Number of sequences in database: 366,079 Lambda K Н 0.711 1.37 1.31 Gapped Lambda H 0.711 1.37 1.31

```
Matrix: blastn matrix:1 -3
Gap Penalties: Existence: 5, Extension: 2
Number of Hits to DB: 2,678,570
Number of Sequences: 38492695
Number of extensions: 2678570
Number of successful extensions: 660781
Number of sequences better than 10.0: 190
Number of HSP's better than 10.0 without gapping: 187
Number of HSP's successfully gapped in prelim test: 3
Number of HSP's that attempted gapping in prelim test: 660488
Number of HSP's gapped (non-prelim): 284
length of query: 233
length of database: 23,466,767,293
effective HSP length: 22
effective length of query: 211
effective length of database: 22,619,928,003
effective search space: 4772804808633
effective search space used: 4772804808633
T: 0
A: 0
X1: 6 (11.9 bits)
X2: 15 (29.7 bits)
S1: 12 (24.3 bits)
S2: 20 (40.1 bits)
</PRE>
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